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## SEQUENCE LISTING

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## 1008889 OSEGUE

548

Lys. Ala Leu Pro Arg Ser 160 accgctgccg gtggtaacca gtggaagacc ccagccccca gggagaggac cccgttctat 654 ccccagccat gataataaag ctgctctccc agctgcctct c 695 <210> 13 <211> 1451 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (105)..(1436) <400> 13 actgoctgga aacgggctgg gcctgcctcg gacgccgccg gtgtcgcgga ttctctttcc 60 gcccgctcca tggcggtgga tgcctgactg gaagcccgag tggg atg cgg ctg acg Met Arg Leu Thr cgg aag cgg ctc tgc tcg ttt ctt atc gcc ctg tac tgc cta ttc tcc Arg Lys Arg Leu Cys Ser Phe Leu Ile Ala Leu Tyr Cys Leu Phe Ser 10 ctc tac gct gcc tac cac gtc ttc ttc ggg cgc cgc cgc cag gcg ccg 212 Leu Tyr Ala Ala Tyr His Val Phe Phe Gly Arg Arg Gln Ala Pro 260 gee ggg tee eeg egg gge etc agg aag ggg geg gee eee geg egg gag Ala Gly Ser Pro Arg Gly Leu Arg Lys Gly Ala Ala Pro Ala Arg Glu 40 308 aga cgc ggc cga gaa cag tcc act ttg gaa agt gaa gaa tgg aat cct Arg Arg Gly Arg Glu Gln Ser Thr Leu Glu Ser Glu Glu Trp Asn Pro 55 356 tgg gaa gga gat gaa aaa aat gag caa caa cac aga ttt aaa act agc Trp Glu Gly Asp Glu Lys Asn Glu Gln Gln His Arg Phe Lys Thr Ser ctt caa ata tta gat aaa tcc acg aaa gga aaa aca gat ctc agt gta 404 Leu Gln Ile Leu Asp Lys Ser Thr Lys Gly Lys Thr Asp Leu Ser Val caa atc tgg ggc aaa gct gcc att ggc ttg tat ctc tgg gag cat att 452 Gln Ile Trp Gly Lys Ala Ala Ile Gly Leu Tyr Leu Trp Glu His Ile ttt gaa ggc tta ctt gat ccc agc gat gtg act gct caa tgg aga gaa 500 Phe Glu Gly Leu Leu Asp Pro Ser Asp Val Thr Ala Gln Trp Arg Glu 125

gga aag tca atc gta gga aga aca cag tac agc ttc atc act ggt cca

		•														
Gly.	Lys	Ser 135	Ile	Val	Gly	Arg	Thr 140	Gln	Tyr	Ser	Phe	Ile 145	Thr	Gly	Pro	
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gct Ala	gtt Val	gtt Val	ttg Leu 200	ctc Leu	gga Gly	aat Asn	gaa Glu	cat His 205	tgt Cys	gat Asp	aat Asn	gag Glu	tgg Trp 210	ata Ile	aac Asn	740
cca Pro	ttc Phe	ctc Leu 215	aaa Lys	aga Arg	aat Asn	gga Gly	ggc Gly 220	ttc Phe	gtg Val	gag Glu	ctg Leu	ctt Leu 225	ttc Phe	ata Ile	ata Ile	788
tat Tyr	gac Asp 230	agc Ser	ccc Pro	tgg Trp	att Ile	aat Asn 235	gac Asp	gtg Val	gat Asp	gtt Val	ttt Phe 240	cag Gln	tgg Trp	cct Pro	tta Leu	836
gga Gly 245	gta Val	gca Ala	aca Thr	tac Tyr	agg Arg 250	aat Asn	ttt Phe	cct Pro	gtg Val	gtg Val 255	gag Glu	gca Ala	agt Ser	tgg Trp	tca Ser 260	884
atg Met	ctg Leu	cat His	gat Asp	gag Glu 265	agg Arg	cca Pro	tat Tyr	tta Leu	tgt Cys 270	aat Asn	ttc Phe	tta Leu	gga Gly	acg Thr 275	att Ile	932
tat Tyr	gaa Glu	aat Asn	tca Ser 280	tcc Ser	aga Arg	cag Gln	gca Ala	cta Leu 285	atg Met	aac Asn	att Ile	ttg Leu	aaa Lys 290	aaa Lys	gat Asp	980
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							aag Lys									1076
							gtc Val									1124
atc Ile	tat Tyr	gag Glu	gct Ala	tgc Cys 345	tcc Ser	tat Tyr	ggc Gly	tcc Ser	att Ile 350	cct Pro	gtg Val	gtg Val	gaa Glu	gac Asp 355	gtg Val	1172
atg Met	aca Thr	gct Ala	ggc Gly	aac Asn	tgt Cys	Gly ggg	aat Asn	aca Thr	tct Ser	gtg Val	cac His	cac His	ggt Gly	gct Ala	cct Pro	1220

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			ctc			atg Met		gct					atc			1268
	-	-			-	gtt Val 395		-							tta Leu	1316
	-			-	-	aga Arg		-			-					1364
	_					atg Met							-	-		1412
		-	aat Asn 440			agt Ser	taa	ttat	cttt	tt g	gagct	=				1451

Alle